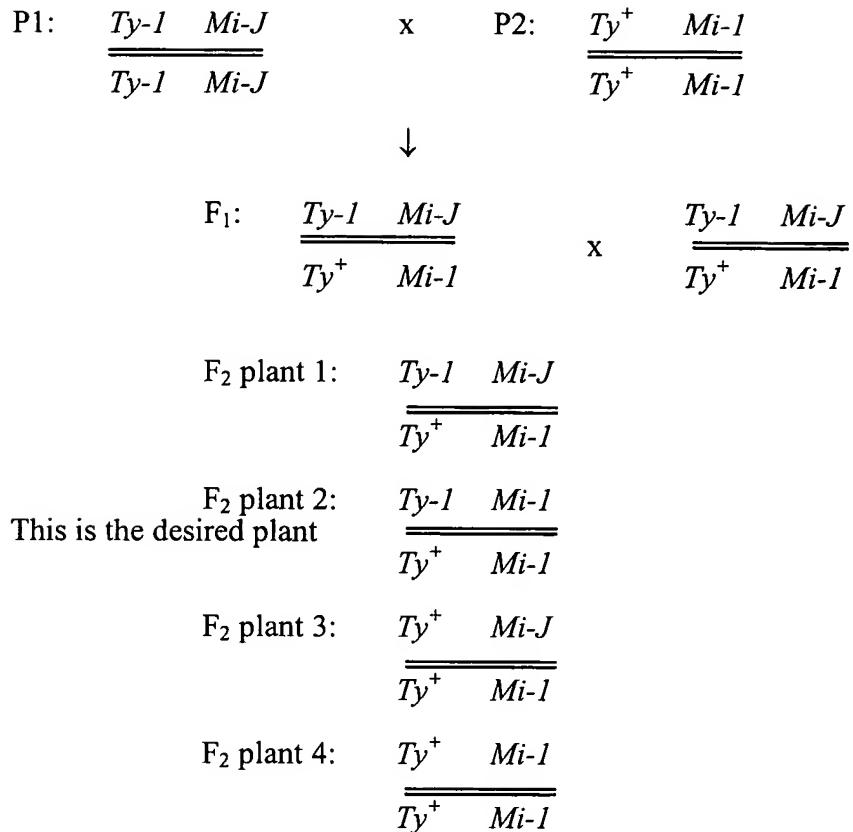


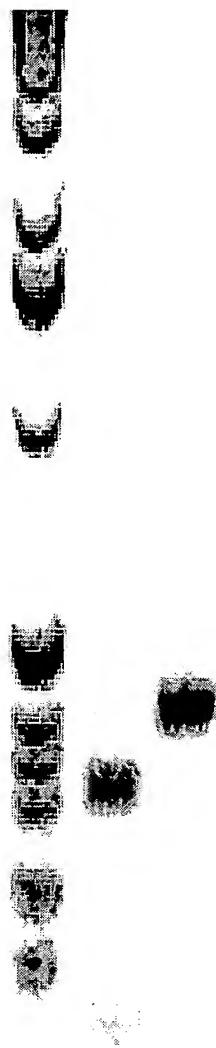
FIGURE 1



This is the desired plant
 as well as other genotypes would be produced in the F2. By selfing plant number 2 completely homozygous plants for $Ty-1$ and $Mi-1$ can be identified.

FIGURE 2

A B



Ty-1 assay

A = resistant allele, 303 & 95

bp

B = susceptible allele, 398 bp

FIGURE 3

A B

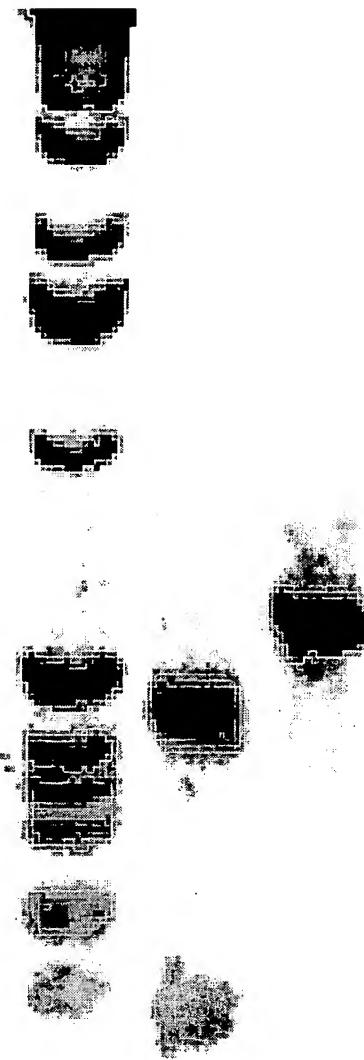


FIGURE 4

A B

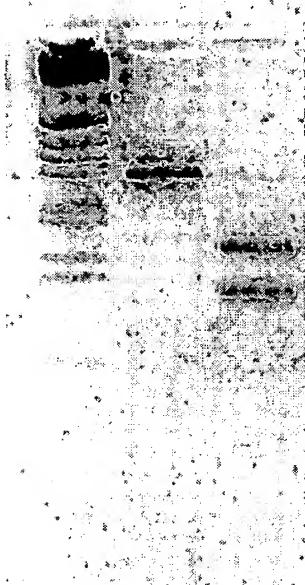


FIGURE 5

$T_y^-/-$ SEQ ID NO.: 10
 $T_y^+/-$ SEQ ID NO.: 11

SEQ ID NO:1

5' TAAATCCGTCGT TACCTCTCC TT

1 C T A A T C C C G T C G T A C C T C C C T T G A A C T A A A A T T T T G T C A A A A G T $T_y^-/-$
1 C T A A T C C C G T C G T A C C T C C C T T G A A C T A A A A T T T G T C A A A A G T $T_y^+/-$

51 A C A A A T C T G T T A T T T A T A T T T T T C T T G G A A T A C T A T C G A T A $T_y^-/-$
51 A C A A A T C T G T T A T T T A T A T T T T C T T G G A A T A C T A T C G A T A $T_y^+/-$

101 T T T T G T A A T A G A A G G T T A G A A T T G T G A T T A T A T G T G A T T G G A A C $T_y^-/-$
100 T T T T G T A A T A G A A G G T T A G A A T T G T G A T T A T A T G T G A T T G G A A C $T_y^+/-$

151 A T T T G T T G C C T T A T G G T G C A A T T A T G T T A C A T G T G T C A T T G G C $T_y^-/-$
150 A T T G T T G C C T T A T G G T G C A A T T A T G T T A C A T G T G T C A T T G G C $T_y^+/-$

201 A C T T A C T G A G T C A T C T T A C T T T A A T A A G A A T T G C T C A A A T G T T T A $T_y^-/-$
200 A C T T A C T G A G T C A T C T T A C T T T A A T A A G A A T T G C T C A A A T G T T T A $T_y^+/-$

251 A A T T C A T T A G C T C A A T G G T A A T T G T A A T T G C A T A T A T C T T T T $T_y^-/-$
250 A A T T G C T T A G C T C A A T G G T A A T T G T A A T T G C A T A T A T C T T T T $T_y^+/-$

301 T G T T C T A G T T C T G A T T A T C A T G T A N C G A A A C T T A T A A A A A T A A $T_y^-/-$
300 T G T T C T A G T T C T G A T T A T C A T G T A C G G A A A C T T A T A A A A A T A A $T_y^+/-$

351 T A G T A A T A G T A G A A N A T T A T G A C A T C A T G C T A T T G A A G T C A T C C $T_y^-/-$
350 T A G T A A T A G T A G T A G A A T T A T G A C A T C A T G C T A T T G A A G T C A T C C $T_y^+/-$

401 G A A T C T A W C
399 G A A T C T A W C

SEQ ID NO:2

FIGURE 6

Mi+ - SEQ ID NO.: 7
Mi-1 - SEQ ID NO.: 8
Mi-J - SEQ ID NO.: 9

FIGURE 7

Mi+ (SEQ ID. NO. 7)

GACACGGACCCACTATTCTGAAACTGATGGTCATTCTTCTCCTTATCGGAGCCTGGCTGAGTTCCAGTCTTGCA
AGCAAAGTGAAGCTGACGTAAGGGATCTGCACTTACATCGTATCCTGTTGAGTTGCAATAACCAGAAACCATGGACT
TTGCTTGAACCTTTTACCTGATTACGATGAACATCTTCTCTAATTCACTGTTCAAGATAATAGATCATAACTCTTG
CCATTGCAGGCACTTACCTCTTAACCATACTGGATTATTGAGAACACATCATTTCACCATCAGAAGACCTTGGG
ACTAGAAGTGGCTAAGGCTGAAGAGGGAGCAACAGAAGGGTGCAGAATTGCAAGATCCTTGTGAAGAATCTGCAGCTT
TAACACTCAACAAAGATAGAGTACTATCCAGATCTGCGGAACTTCACTGTTTAACTTGACCTGTTCCAGCACTA
CCATTGCTGCACTAGTGTCTCCGGTCAAGAACGGAGACCTGCTACCTTCTCTAGAGATGTCATCACATAT
TTTTCCATAGAATCTGGGATTACATGTCAGGAATCTCGCAGTCTCTCCCTTCTCTTAATCGGAGAACAT
TGTCAACTTCCCTTATGCGTTGACACATCGGAAATATAAGCTCTGGGTTTGTGAACCTCATCTAGACTTT
TCATCCTCTGAGTCCCTGTTTACATTGTCAGGAATCATCTGGCATTACTGCTGAACCTCATCTAGACTTT
AACACAGGGCAAAGGCTGGTCTCGTCATCGAGTCATCATCTGTATAATTGGAGAACATCTGATTCCA
CTTCACTTGTGTTCTTCTATTGACATCCCGTAG

Mi-1 (SEQ ID. NO. 8)

GACACGGACCCACTATTCTGAAACCGATGGTCATTCTTCTCCTTATCGGAGCCTGGCTGAAATTCCCGTCTTGCA
AGCAAATTGAAGCTGACGTAAGGGATCTGCACTTGCATCGTATCCTGTTGAGTTGCAATAACCAGAAACCGTGGACT
TTGCTTGAACCTTTTACCTGATTACGATGGACACTTCTCTCTAATTCACTGTTCAAGATAATAGATCATAACTCTTG
CCATTGCAGGCACTTACCTCTTAACCATACTGGATTATTGAGAACACATCATTTCACCATCAGAAGACCTTGGG
ACTAGAAGTGGGAAAGGCTGAAGAGGGAGCAACAGAAGGGTGCAGAATTGCAAGATCCTTGTGAAGAATCTGCAGCTT
TAACACTCAACAAAGATAGAGTACTATCCAGATCTGCCCAGCCTGCTGTTCTTTAACTTGACCTGTTCCAGCACTA
CCATTGCTGCACTAGTGTCTCCGGTCAAGAACGGAGACCTGCTACCTTCTCTGGAGATGTCATCACATAT
TTTTCCATAGAATCTGGGATTACATGTCAGGAATCTCGAAGTCTCTCCCTTCTCTTAATCGGAGAACAT
TGTCAACTTCCCTTATGCGTTGACACATCGGAAATATAAGCTCTGGGTTTGTGAACCTCATCTAGACTTT
TCATCCTCTGAGTCCCTGTTTACATTGTCAGGAATCATCTGGCATTACTGCTGAACCTCATCTAGACTTT
AACACAGGGCAGAAGGCTGGTCTCGTCATCGAGTCATCATCTGTATAATTGGAGAACATCTGATTCCA
CTTCACTTGTGTTCTTCTATTGACATCCCGTAG

Mi-J (SEQ ID. NO. 9)

GACACGGACCCACTATTCTGAAACTGATGGTCATTCTTCTCCTTATCGGAGCCTGGCTGACTTTCCAGTCTTGCA
AGCAAATTGAAGCTGACGTAAGGGATCTGCACTTACATCGTATCCTGTTGAGTTGCAATAACCAGAAACCGTGGACT
TTGCTTGAACCTTTTACCTGATTACGATGGACACTTCTCTCTAATTCACTGTTCAAGATAATAGATCATAACTCTTG
CCATTGCAGGCACTTACCTCTTAACCATACTGGATTATTGAGAACACATCATTTCACCATCAGAAGACCTTGGG
ACTAGAAGTGGTAAGGCTGAAGAGGGAGCAACAGAAGGGTGCAGAATTGCAAGATCCTTGTGAAGAATCTGCAGCTT
TAACACTCAACAAAGATAGAGTACTATCCAGATCTGCCCAGCCTGCTGTTCTTTAACTTGACCTGTTCCAGCACTA
CCATTGCTGCACTAGTGTCTCCGGTCAAGAACGGAGACCTGCTACCTTCTCTGGAGATGTCATCACATAT
TTTTCCATAGAATCTGGGATTACATGTCAGGAATCTCGAAGTCTCTCCCTTCTCTTAATCGGAGAACAT
TGTCAACTTCCCTTATGCGTTGACACATCGGAAATATAAGCTCTGGGTTTGTGAACCTCATCTAGACTTT
TCATCCTCTGAGTCCCTGTTTACATTGTCATGAATCATCTGGCATTACTGCTGAACCTCATCTAGACTTT
AACACAGGGCAGAAGGCTGGTCTCGTCATCGAGTCATCATCTGTATAATTGGAGAACATCTGATTCCA
CTTCACTTGTGTTCTTCTATTGACATCCCGTAG

Ty-1 (SEQ ID. NO. 10)

CTAATCCGCGTTACCTCTCCTTGAACAAAAATTGGTCAAAAGTTACAAATCTGTTATTATATATTTTT
CTTGGAAATTACTATCGATAATTGGTAAATTAGAAGGTTAGAATTGGAGTATATGTTGTGATTGGAACGATTGGTTGT
GCCTTATGGTGGAAATTATGTTACATGTGTCATTGGCTAACTTACTGAGTCATCTTACTTTAAATAAGAACATGCTTC
AAATGTTTATAATTCTTAGCTCAATGGTAATTGTTATTGATGCAATATCTTTTGTTCTAGTTCTGATTATA
TCATGTANGAAACTTATATAAAAATAATTGTAATAGTAGTGAANATTGACATATTGCTATTGAAGTCATCCG
GAATCT

Ty + (SEQ ID. NO. 11)

CTAATCCGCGTTACCTCTCCTTGAACAAAAATTGGTCAAAAGTTACAAATCTGTTATTATATATTTTT
TTGGAAATTACTATCTTATTGGTAAATTAGAAGGTTAGAATTGGAGTATATGTTGTGATTGGAACGATTGGTTGT
CCTTATGGTGGAAATTATGTTACATGTGTCATTGGTAACTTACTGAGTCATCTTACTTTAAATAAGAACATGCTTC
TATGTTTATAATTCTTAGCTCAATGGTATTGTTATTGATGCAATATCTTTTGTTCTAGTTCTGATTATA
CATGTAGGAAACTTATATAAAAATAATTGTAATAGTAGTGAANATTGACATATTGCTATTGAAGTCATCCG
ATCTANC

Figure 8 - Normalized Pathology Scoring

